

SEQUENCE LISTING

```
<110>
       Efendic, Suad
<120>
       USE OF GLP-1 OR ANALOGS IN TREATMENT OF MYOCARDIAL INFARCTION
<130>
       X-10822A
<140>
       US 09/834,229
<141>
       2001-04-12
<150>
      US 08/915,918
<151>
       1997-08-21
<150>
       US 06/024,980
<151>
       1996-08-30
<160>
<170> PatentIn version 3.1
<210>
<211>
       31
<212> PRT
<213>
       Homo sapiens
<400> 1
His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
                5
                                     10
Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Gly
                                                     30
<210>
       2
<211>
       31
<212>
       PRT
<213>
      Artificial Sequence
<220>
<223>
       synthetic construct
<220>
<221>
      MISC_FEATURE
<222>
      (1)..(1)
<223>
       Xaa at position 1 is L-histidine, D-histidine, desamino-histidine
       , 2-amino-histidine, B-hydroxy-histidine, homohistidine, alpha-fl
       uoromethyl-histidine, and aplpha-methyl-histidine;
<220>
<221>
      MISC_FEATURE
<222>
       (2)..(2)
<223>
      Xaa at position 2 is Ala, Gly, Val, Thr, Ile, and alpha-methyl-Al
<220>
<221>
      MISC_FEATURE
<222>
       (15)..(15)
<223> Xaa at position 15 is Glu, Gln, Ala, Thr, Ser, and Gly;
```

Page 1





```
<220>
<221> MISC_FEATURE
<222>
      (21)..(21)
<223> Xaa at position 21 is Glu, Gln, Ala, Thr, Ser, and Gly;
<220>
<221> MOD_RES
<222> (31)..(31)
<223> AMIDATION
<220>
<221> MISC_FEATURE
<222> (31)..(31)
<223> Xaa at position 31 is Gly
<400> 2
Xaa Xaa Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Xaa Gly
Gln Ala Ala Lys Xaa Phe Ile Ala Trp Leu Val Lys Gly Arg Xaa
                                25
<210> 3
<211> 29
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<220>
<221> MISC_FEATURE
<222>
       (29)..(29)
<223> Xaa at position 29 is absent or Gly.
<400> 3
His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Xaa
            20
<210>
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<220>
<221> MISC_FEATURE
<222> (19)..(19)
```



```
<223> Xaa at position 19 is Lys or Arg;
<220>
<221> MOD_RES
<222> (30)..(30)
<223> AMIDATION
<220>
<221> MISC_FEATURE
<222> (30)..(30)
<223> Xaa at position 30 is Gly.
<400> 4
Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly Gln
Ala Ala Xaa Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Xaa
<210> 5
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 5
His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
                                    10
Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
<210> 6
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 6
Ser Arg Arg Gln
```